

NT1p MGLFANSTKDFLAGVAAMVSKTAVAPIERVKLLIQVQHASKQITAEKQ 50
 NT2p MITANSTFAKDFLAGGVAALSKTAVAPIERVKLLIQVQHASKQITADKQ 50
 NT3p MITQATSTFAKDFLAGGTAAALSKTAVAPIERVKLLIQVQHASKQITADKQ 50
 NT1p YKGIIDCVVRIPKEQCFLSFWRGNLANVIRYFPTQALNFAFKDKYKQIFL 100
 NT2p YKGIIDCVVRIPKEQGVLSFWRGNLANVIRYFPTQALNFAFKDKYKQIFL 100
 NT3p YKGIIDCVVRIPKEQGVLSFWRGNLANVIRYFPTQALNFAFKDKYKQIFL 100
 NT1p GGVDKHTQFWRYFAGNLASGGAAGATSLCFVYPLDFARTRLAADVGKQGA 149
 NT2p GGVDKHTQFWRYFAGNLASGGAAGATSLCFVYPLDFARTRLAADVGKQGA 150
 NT3p GGVDKHTQFWRYFAGNLASGGAAGATSLCFVYPLDFARTRLAADVGKQGA 150
 NT1p EREFRGLGDCIKKFKSDGIRGLYQGFVSVQGIITZRAAYFGVYDTAKG 199
 NT2p EREFRGLGDCIKKFKSDGIRGLYQGFVSVQGIITZRAAYFGVYDTAKG 200
 NT3p EREFRGLGDCIKKFKSDGIRGLYQGFVSVQGIITZRAAYFGVYDTAKG 200
 NT1p MLPDPKNTIIVSWMTAQSVTAVAGLISYPFDIVRRRMMQSGRKGADIM 249
 NT2p MLPDPKNTIIVSWMTAQSVTAVAGLISYPFDIVRRRMMQSGRKGADIM 250
 NT3p MLPDPKNTIIVSWMTAQSVTAVAGLISYPFDIVRRRMMQSGRKGADIM 250
 NT1p YTGIVDCNRKIAFDEGCKAFFKCAWSNVLRGMGCAFVLVLYDEIKKVV. 298
 NT2p YTGIVDCNRKIAFDEGCKAFFKCAWSNVLRGMGCAFVLVLYDEIKKVV. 299
 NT3p YTGIVDCNRKIAFDEGCKAFFKCAWSNVLRGMGCAFVLVLYDEIKKVV. 299

Figure 2

250 —
148 —
60 —
42 —
30 —
22 —

M 0 G/n 1 2 3 4 5 M

Figure 3

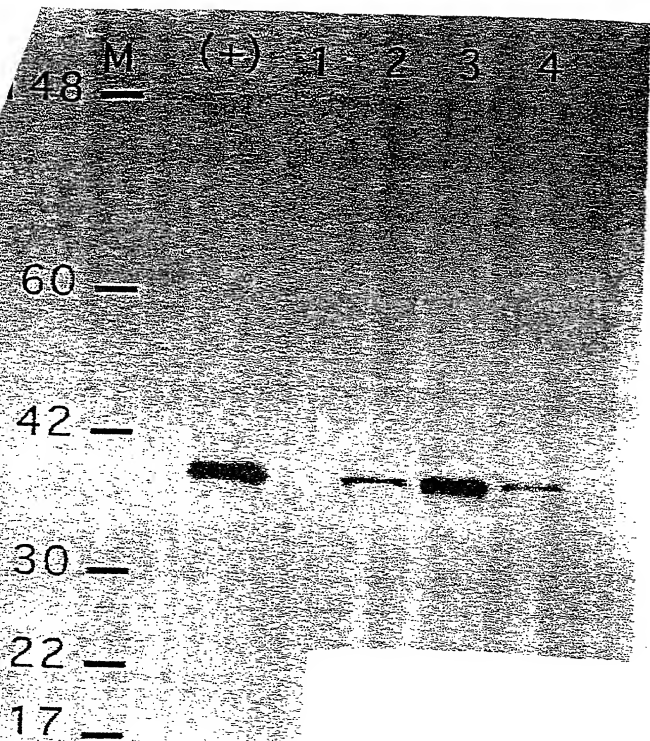


Figure 4

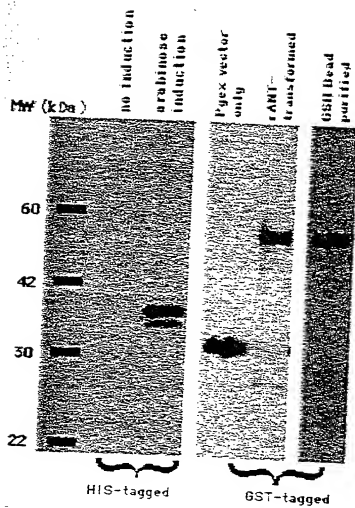


Figure 5

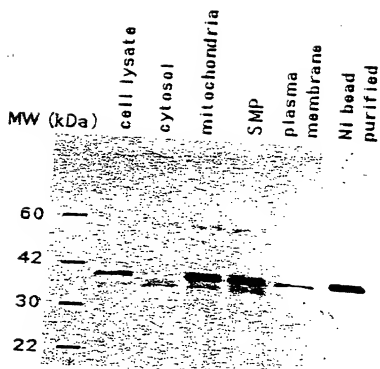


Figure 6

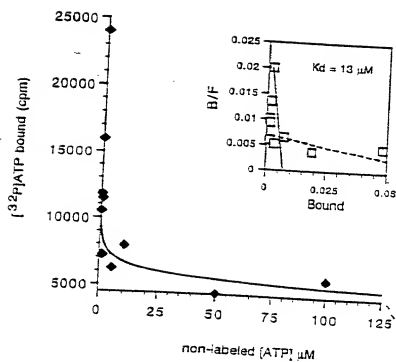


Figure 7

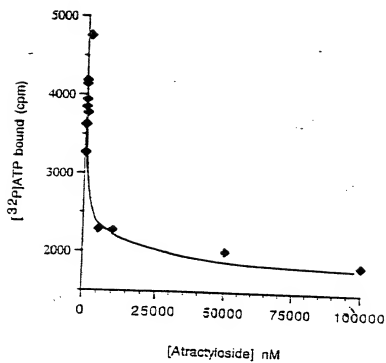


Figure 8

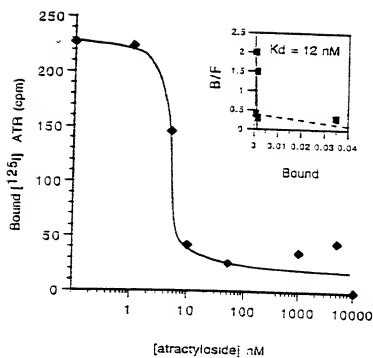


Figure 9

0011131-024001

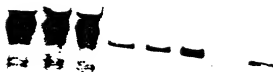


Figure 10 -

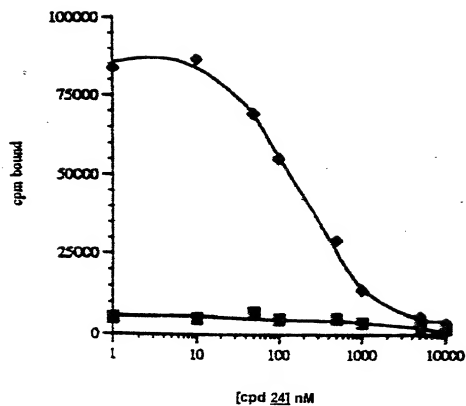


Figure 11

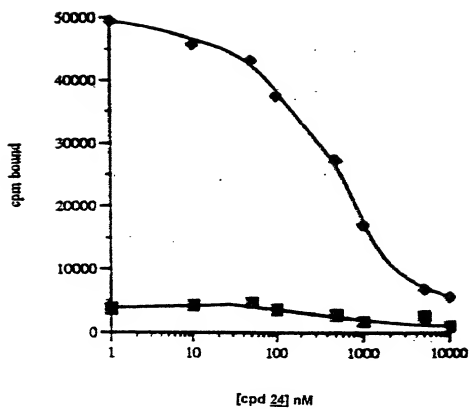


Figure 12

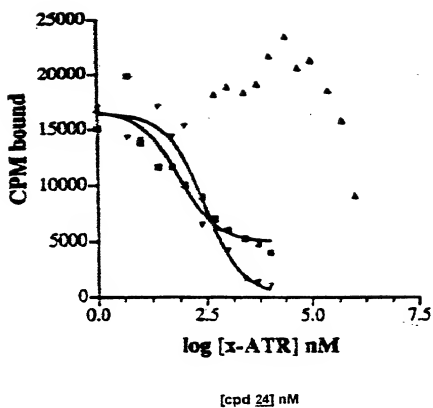


Figure 13

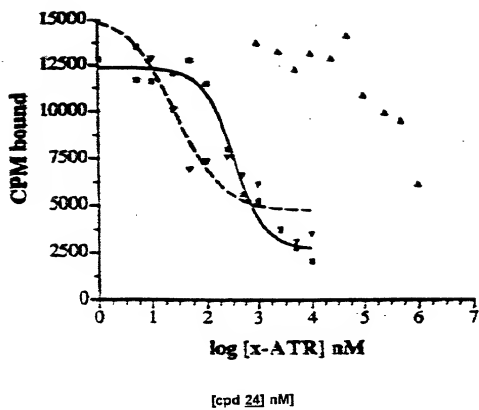


Figure 14

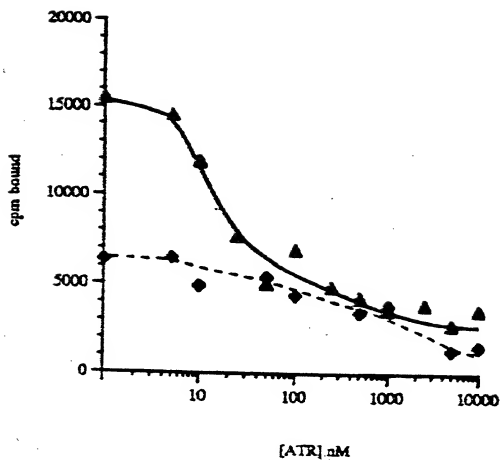


Figure 15

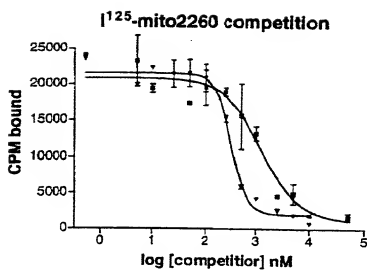


Figure 16

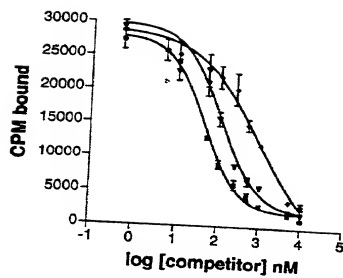


Figure 17

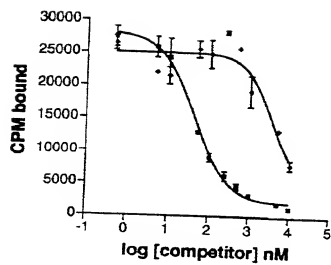


Figure 18

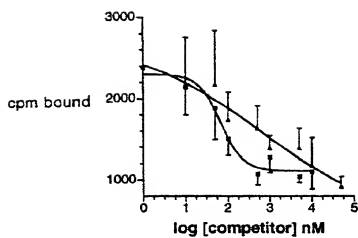


Figure 19